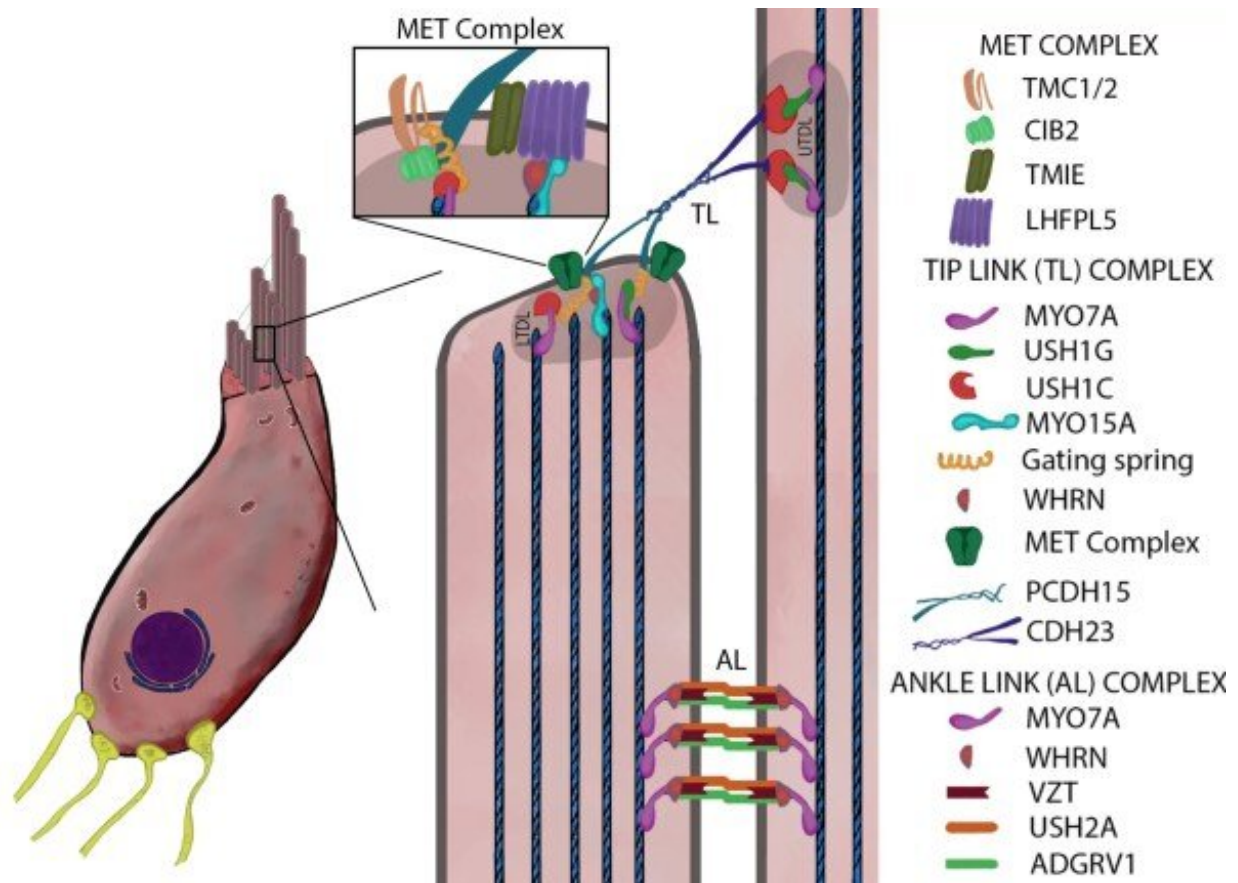


Identification of new genes associated with familial Meniere's disease

December 1 2021



Proteins involved in bundling together the stereocilia of the sensory cells of the inner ear. Credit: University of Granada

A team of scientists from the University of Granada (UGR) and

GENYO (Pfizer-University of Granada-Andalusian Government Centre for Genomics and Oncological Research—led by Pablo Roman-Naranjo and Jose Antonio López-Escámez, researcher in charge of the Biohealth Research Institute in Granada (ibs. GRANADA)—has identified new genes associated with familial Meniere's disease.

Meniere's disease is a disorder of the inner ear that is characterized by repeated episodes of vertigo and persistent noise in the ears (tinnitus), leading to progressive hearing loss.

This disease has a hereditary component, as identified in family aggregation studies, with approximately 10% of patients having one or more relatives affected by it. In this new study, published in the journal *Hearing Research*, a total of 62 families affected by this disease (mainly from Spain) were analyzed. The results point to the existence of a relationship between familial Meniere's disease and various ear-[genes](#), the main candidate being the MYO7A gene, which encodes a protein called myosin VIIa that is specific to inner-ear hair cells.

The correct functioning of the ear relies on thousands of genes. When these genes are altered by rare [mutations](#), this may reduce both the individual's ability to keep their balance and to hear properly. In the case of Meniere's disease, the researchers in this study propose a "digenic inheritance" model, in which rare mutations in at least two genes are necessary.

The origin of this disease in some families seems to lie in rare mutations of the MYO7A gene, together with mutations in other genes, such as CDH23, PCDH15, or ADGRV1 (that is, genes with which MYO7A interacts). This hypothesis is based on the results obtained in the present study: nine of the families presented rare mutations in these genes.

As the article notes, these genes are expressed in a specific area of the ear: the stereocilia of the neurosensory cells of the inner ear. These cells are responsible for the transmission of sound in the organ of Corti, as well as the perception of acceleration in the posterior labyrinth, sending [nerve impulses](#) to the central nervous system. Stereocilia are found in each of these cells. According to the authors, mutations in MYO7A and other genes would cause alterations in several proteins that keep the stereocilia bundled together, ultimately causing hearing loss and vertigo.

"We suggest that [rare mutations](#) in the MYO7A gene, either by itself or combined with other mutations in genes that interact with MYO7A, would cause alterations in the morphostructure of the stereocilia and a loss of cohesiveness between them, with an abnormal opening of the mechanotransduction complex (located in the cilia) and, finally, hearing loss and/or vertigo. The MYO7A gene has been associated with familial sensorineural hearing loss or [retinitis pigmentosa](#)—another rare disease that causes blindness," explain the authors.

The study of Meniere's [disease](#) and the identification of [new genes](#) that can explain its origin can help to decipher the mechanisms underlying it and thereby facilitate its early genetic diagnosis, as well as contribute to the development and application of new drugs in its treatment.

More information: P. Roman-Naranjo et al, Rare coding variants involving MYO7A and other genes encoding stereocilia link proteins in familial meniere disease, *Hearing Research* (2021). [DOI: 10.1016/j.heares.2021.108329](#)

Provided by University of Granada

Citation: Identification of new genes associated with familial Meniere's disease (2021, December

1) retrieved 25 April 2024 from

<https://medicalxpress.com/news/2021-12-identification-genes-familial-meniére-disease.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.